SIN

(FILE 'HOME' ENTERED AT 14:25:29 ON 07 JAN 2008)

L14

W

	(FILE HOME, ENTERED AT 14:25:29 ON 07 DAM 2008)
L1	FILE 'REGISTRY' ENTERED AT 14:25:37 ON 07 JAN 2008 296092 S RKR/SQSP
L2 L3 L4	
L5	FILE 'REGISTRY' ENTERED AT 14:28:47 ON 07 JAN 2008 0 S RKR/SPSP AND PEG
L6 L7 L8 L9 L10 L11	FILE 'CAPLUS, MEDLINE' ENTERED AT 15:12:37 ON 07 JAN 2008 338 S BRANCHED (P) (LYS OR LYSINE) (P) (ARG OR ARGININE) 4 S L6 (P) RESIN 2 S L6 (P) IMMOBILIZ? 12 S L6 (P) AFFINITY 18 S L7 OR L8 OR L9 24 S ARG (A) LYS (A) 2 1 S L11 AND (RESIN OR AFFINITY OR IMMOBILIZ? OR DENDRIMER)
L13	FILE 'CAPLUS, MEDLINE' ENTERED AT 15:20:13 ON 07 JAN 2008 35 S (LYS OR LYSINE) (A) (ARGININE) (A) 2

25 S L13 AND (RESIN OR AFFINITY OR IMMOBILIZ? OR PEPTIDE)

(FILE 'HOME' ENTERED AT 09:54:11 ON 07 JAN 2008)

FILE 'REGISTRY' ENTERED AT 09:54:22 ON 07 JAN 2008 L1 5 S RKR/SQEP

FILE 'CAPLUS, BIOSIS' ENTERED AT 09:54:41 ON 07 JAN 2008 L2 3 S L1

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	FILE 'CAPLU	JS	, BIOSIS' ENTERED AT 09:59:47 ON 07 JAN 2008
L3	166982	S	POLYSTYRENE
L4	165846	S	(POLYETHYLENE (A) GLYCOL) OR PEG
L5	20171	S	TRIPEPTIDE
L6	2	S	L3 (P) L4 (P) L5
L7	4	S	L3 AND L4 AND L5
L8	2	S	L7 NOT L6
L9	259332	S	(ARGININE OR ARG)
L10	34	S	L9 AND L3 AND L4
L11	24	S	L10 AND (RESIN OR IMMOBILIZ? OR AFFINITY)
L12	20	S	L11 AND PEPTIDE
L13	1632	S	L9 AND (PEPTIDE (P) COLUMN)
L14	354	S	L13 AND AFFINITY
L15	343	S	L9 AND (PEPTIDE (P) COLUMN (P) AFFINITY)
L16	121	S	L15 AND (LYS OR LYSINE)
L17	0	S	L16 AND L3 AND L4
L18	7	S	L16 AND RESIN
L19	415	S	RAPP W?/AU
L20	6	S	L19 AND AFFINITY AND POLYSTYRENE
L21	173	S	DEPPISCH R?/AU
L22		-	L21 AND AFFINITY AND POLYSTYRENE NOT L20
L23	0	S	GOHL H?/AU AND AFFINITY AND POLYSTYRENE NOT L20
L24			WITTNER B?/AU
L25			WITTNER B?/AU AND AFFINITYA AND POLYSTYRENE NOT L20
L26			WITTNER B?/AU AND AFFINITY AND POLYSTYRENE NOT L20
L27	1	S	BECK W?/AU AND AFFINITY AND POLYSTYRENE NOT L20

FILE 'REGISTRY' ENTERED AT 16:05:58 ON 07 JAN 2008 L15 STRUCTURE UPLOADED

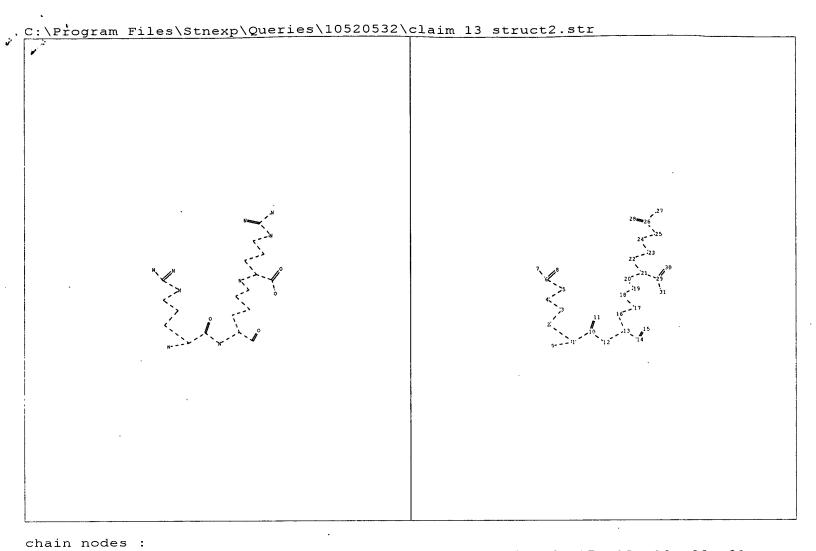
FILE 'CAPLUS' ENTERED AT 16:06:37 ON 07 JAN 2008

FILE 'REGISTRY' ENTERED AT 16:06:51 ON 07 JAN 2008

0 S L15 SSS SAM

L17 0 S L15 SSS FULL

L16



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1 2 3 4 5 6 7 8 9 10 11 12 13 14 15 16 17 18 19 20 21 22 23 24 25 26 27 28 29 30 31  

chain bonds:
    1-2 1-9 1-10 2-3 3-4 4-5 5-6 6-7 6-8 10-11 10-12 12-13 13-14 13-16 14-15 16-17 17-18 18-19 19-20 20-21 21-22 21-29 22-23 23-24 24-25 25-26 26-27 26-28 29-30 29-31  

exact/norm bonds:
    1-2 1-9 1-10 2-3 3-4 4-5 5-6 6-7 6-8 10-11 10-12 12-13 13-14 13-16 14-15 16-17 17-18 18-19 19-20 20-21 21-22 21-29 22-23 23-24 24-25 25-26 26-27 26-28 29-30 29-31  

Match level:
    1:CLASS 2:CLASS 3:CLASS 4:CLASS 5:CLASS 6:CLASS 7:CLASS 8:CLASS
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9:CLASS 10:CLASS 11:CLASS 12:CLASS 13:CLASS 14:CLASS 15:CLASS 16:CLASS 17:CLASS 18:CLASS 19:CLASS 20:CLASS 21:CLASS 22:CLASS 23:CLASS 24:CLASS 25:CLASS 26:CLASS 27:CLASS 28:CLASS 29:CLASS

23:CLASS 24:CLASS 30:CLASS 31:CLASS